

FORM PTO-1449 U.S. DEPARTMENT OF COMMERCE PATENT AND TRADEMARK OFFICE		ATTY. DOCKET NO. NADII.030A	APPLICATION NO. 09/938,714
INFORMATION DISCLOSURE STATEMENT BY APPLICANT		APPLICANT Chantal Roth	
(USE SEVERAL SHEETS IF NECESSARY)		FILING DATE August 23, 2001	GROUP 2175

## U.S. PATENT DOCUMENTS

## FOREIGN PATENT DOCUMENTS

EXAMINER INITIAL	DOCUMENT NUMBER	DATE	COUNTRY	CLASS	SUBCLASS	TRANSLATION	
						YES	NO
<i>DR</i>	6 WO 99/12040	03/11/99	PCT			<i>✓</i>	
<i>DR</i>	7 WO 99/61654A1	12/02/99	PCT			<i>✓</i>	
<i>DR</i>	8 WO 00/11206A1	03/02/00	PCT			<b>RECEIVED</b>	
<i>DR</i>	9 WO 00/29987	05/25/00	PCT			<i>✓</i>	
						<i>DEC 30 2002</i>	
						<b>Technology Center 2100</b>	

**EXAMINER  
INITIAL**

**OTHER DOCUMENTS (INCLUDING AUTHOR, TITLE, DATE, PERTINENT PAGES, ETC.)**

INITIAL

 10 Washburn et al., Large-scale analysis of the yeast proteome by multidimensional protein identification technology, *Nature Biotechnology* 19:242-247 (2001)

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EXAMINER 	DATE CONSIDERED <u>3/21/04</u>
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1. Conrads, et al., Quantitative Analysis of Bacterial and Mammalian Proteomes Using a Combination of Cysteine Affinity Tags and <sup>15</sup>N-Metabolic Labeling; *Anal. Chem.* 73:2132-2139 (2001)
2. Griffin et al., Quantitative Proteomic Analysis Using a MALDI Quadrupole Time-of-Flight Mass Spectrometer; *Anal. Chem.* 73:978-986 (2001)
3. Gygi, et al., Quantitative analysis of complex protein mixtures using isotope-coded affinity tags; *Nature Biotech* 17:994-999; (1999)
4. Han et al., Quantitative profiling of differentiation-induced microsomal proteins using isotope-coded affinity tags and mass spectrometry; *Nature Biotech* 19:946-951 (2001)
5. Oda, et al., Accurate quantitation of protein expression and site-specific phosphorylation; *Proc. Natl. Acad. Sci. USA* 96:6591-6596 (1999)
6. Yao, et al., Proteolytic <sup>18</sup>O Labeling for Comparative Proteomics: Model Studies with Two Serotypes of Adenovirus; *Anal. Chem.* 73:2836-2842 (2001)

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EXAMINER		DATE CONSIDERED	
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